

1  
SEQUENCE LISTING

<110> FRENKEL LEO G.  
VAN DER LINDEN CORNELIS P.

<120> METHOD FOR PRODUCING ANTIBODY FRAGMENTS

<130> 060113-0271592

<140> 09/626,242

<141> 2000-09-27

<150> PCT/EP99/00481

<151> 1999-01-25

<150> EP 98300525.7

<151> 1998-01-26

<160> 18

<170> PatentIn Ver. 2.1

<210> 1

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 1

aggtsmarct gcagsagtcw gg

22

<210> 2

<211> 57

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Primer

<400> 2

catgccatga ctgcggccc agccggccat ggccsaggtg marctgcags agtcwgg

57

<210> 3

<211> 53

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Primer

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aacagttaag cttccgcttg cggccgcgga gctgggggtct tcgctgtggt gcg

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<210> 4  
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 <212> DNA  
 <213> Artificial Sequence

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 <223> Description of Artificial Sequence: Primer

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<210> 5  
 <211> 117  
 <212> PRT  
 <213> Lama glama

<400> 5  
 Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Asp  
 1 5 10 15  
 Phe Leu Arg Phe Ser Cys Ala Ala Leu Gly Ala Arg Phe Ser Ser Asp  
 20 25 30  
 Val Met Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val  
 35 40 45  
 Ala Ala Ser Ser Trp Asn Gly Asp Thr Thr His Tyr Ser Asp Ser Val  
 50 55 60  
 Glu Gly Gln Phe Thr Ile Ser Arg Asp Ile Ala Lys Asn Thr Ser Tyr  
 65 70 75 80  
 Leu Gln Met Asn Arg Leu Gln Pro Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Arg Trp Cys Arg Pro Pro Arg Pro Lys Tyr Trp Gly Gln Gly Thr Gln  
 100 105 110  
 Val Thr Val Ser Ser  
 115

<210> 6  
 <211> 115  
 <212> PRT  
 <213> Lama glama

<400> 6  
 Gln Val Gln Leu Gln Gln Ser Gly Gly Gly Leu Val Gln Ala Gly Ser  
 1 5 10 15  
 Phe Leu Ser Phe Ser Cys Thr Ala Ser Gly Arg Thr Phe Ser Asn Tyr  
 20 25 30  
 Ala Met Gly Trp Phe Arg Gln Ala Ser Gly Asn Gln Arg Ala Phe Val  
 35 40 45

Ala Ala Ile Gly Arg Asn Gly Asp Thr His Tyr Ile Asp Ser Val Lys  
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Gly Lys Asp Thr Val Tyr Leu  
65 70 75 80

Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys Arg  
85 90 95

Ile Trp Val Gly Ala Arg Asp Tyr Trp Gly Gln Gly Thr Gln Val Thr  
100 105 110

Val Ser Ser  
115

<210> 7

<211> 116

<212> PRT

<213> Lama glama

<400> 7

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Gly  
1 5 10 15

Phe Leu Arg Phe Ser Cys Ala Ala Ser Gly Arg Thr Phe Ser Arg Tyr  
20 25 30

Thr Met Gly Trp Phe Arg Gln Ala Pro Gly Asn Glu Arg Lys Phe Val  
35 40 45

Ala Ala Val Ser Thr Ser Gly Asn Thr His Tyr Thr Gly Ser Val Lys  
50 55 60

Gly Arg Phe Thr Ile Phe Arg Gln Asn Ala Lys Asn Thr Val Tyr Leu  
65 70 75 80

Gln Met Ser Asn Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys Ala  
85 90 95

Ala Arg Phe Gly Gly Met Asn Trp Lys Tyr Trp Gly Gln Gly Ile Gln  
100 105 110

Val Thr Val Ser  
115

<210> 8

<211> 121

<212> PRT

<213> Lama glama

<400> 8

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Pro  
1 5 10 15

Phe Leu Asn Val Ser Cys Val Val Ser Gly Gly Ile Phe Ser Asp Tyr  
20 25 30

Thr Leu Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Lys Phe Val  
35 40 45

Ala Ala Val Ser Ser Gly Gly Ser Thr His Tyr Thr Gly Ser Val Lys  
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Ala Asn Thr Met Tyr Leu  
65 70 75 80

Gln Met Ser Ser Leu Lys Pro Asp Asp Thr Ala Val Tyr Tyr Cys Asn  
85 90 95

Ala Ile Val Pro Pro Thr Arg Thr Phe Cys Gly Arg Thr Tyr Trp Gly  
100 105 110

Gln Gly Thr Gln Val Thr Val Ser Ser  
115 120

<210> 9

<211> 112

<212> PRT

<213> Lama glama

<400> 9

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Asp  
1 5 10 15

Phe Val Arg Leu Ser Cys Ala Ala Ser Arg Arg Ala Ser Ser Thr Tyr  
20 25 30

Ala Val Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val  
35 40 45

Gly Arg Ile His Arg Gly Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Thr Gln Asn Thr Val Tyr  
65 70 75 80

Leu Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Asn Val Arg Ser Tyr Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser  
100 105 110

<210> 10

<211> 117

<212> PRT

<213> Lama glama

<400> 10

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Gly  
1 5 10 15

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<210> 11
<211> 124
<212> PRT
<213> Lama glama
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<400> 11
Gln Val Lys Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Gly
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  20           25           30
Ala Ile Gly Trp Phe Arg Gln Ala Pro Gly Lys Gln Arg Glu Leu Val
  35           40           45
Ala Gly Ile Ser Thr Gly Gly Ser Thr Asn Tyr Ala Asp Ser Val Lys
  50           55           60
Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asp Thr Val Tyr Leu
  65           70           75           80
Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys Ala
  85           90           95
Ala Gly Arg Arg Ile Ser Ser Ser Tyr Tyr Ser Arg Gly Leu Tyr Ala
  100          105          110
Tyr Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser
  115          120

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<210> 12
<211> 124
<212> PRT
<213> Lama glama
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&lt;400&gt; 12

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Asp  
 1 5 10 15

Ser Leu Arg Leu Ser Cys Glu Ala Ser Gly Arg Ser Phe Ser Asn Phe  
 20 25 30

Ala Met Ala Trp Phe Arg Gln Thr Pro Gly Lys Glu Arg Glu Phe Val  
 35 40 45

Ala Gly Ile Ser Trp Arg Gly Gly Arg Thr Tyr Tyr Ala Ala Ser Val  
 50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Gly Lys Asn Thr Val Tyr  
 65 70 75 80

Leu Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95

Ala Thr Ala Tyr Gly Gln Gly Pro Ile Thr Val Pro Lys Phe Tyr Thr  
 100 105 110

Tyr Arg Gly Gln Gly Thr Gln Val Thr Val Ser Ser  
 115 120

&lt;210&gt; 13

&lt;211&gt; 121

&lt;212&gt; PRT

&lt;213&gt; Lama glama

&lt;400&gt; 13

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Gly  
 1 5 10 15

Cys Val Arg Leu Ser Cys Ala Ala Ser Gly Arg Thr Phe Ser Arg Tyr  
 20 25 30

Thr Met Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val  
 35 40 45

Ala Ala Ile Ser Trp Arg Ser Gly Gly Ile Lys Ile Tyr Gly Asp Ser  
 50 55 60

Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asp Thr Val  
 65 70 75 80

Tyr Val Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr  
 85 90 95

Cys Asn Ser Arg Pro Arg Ile Tyr Arg Gly Asn Val Val Tyr Trp Gly  
 100 105 110

Gln Gly Thr Gln Val Thr Val Ser Ser  
 115 120

<210> 14  
 <211> 34  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Synthetic  
 oligonucleotide

<400> 14  
 ggcccagccg gccatggccc aggtgcagct gcag

34

<210> 15  
 <211> 11  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Synthetic  
 peptide

<400> 15  
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 1 5 10

<210> 16  
 <211> 39  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Synthetic  
 oligonucleotide

<400> 16  
 gcggccgccc atcaccatca ccatcacggg gccgcagaa

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 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Synthetic  
 peptide

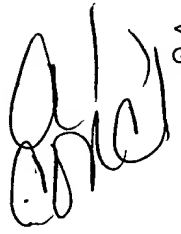
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 Ala Ala Ala His His His His His His Gly Ala Ala Glu  
 1 5 10

<210> 18  
 <211> 11  
 <212> PRT  
 <213> Unknown Organism

&lt;220&gt;

<223> Description of Unknown Organism: Myc peptide  
sequence

&lt;400&gt; 18

Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn  
1 5 10

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